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TECH CENTER

CRF Errors Edited by the STIC Systems Branch

Serial Number: 08/887,505A CRF Edit Date: 9/15/2003
Edited by: Are

ENTERED

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

J ___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

98

___ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

_____

___ Moved responses to same line as heading/numeric identifier, specifically:

_____

___ Other:

_____

_____

_____

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SEP 16 2003
TECH CENTER 1600

RAW SEQUENCE LISTING DATE: 09/15/2003
PATENT APPLICATION: US/08/887,505A TIME: 10:39:53

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\09152003\H887505A.raw

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Kilkuskie, Robert E.
Frank, Bruce L.
Goodchild, John
Wolfe, Jia L.
Roberts, Peter C.
Hamlin, Jr., Henry A.
Roberts, Noel A.
Walther, Debra M.

(ii) TITLE OF INVENTION: Oligonucleotides Specific for Hepatitis C Virus

(iii) NUMBER OF SEQUENCES: 173

(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Hale and Dorr LLP
(B) STREET: 60 State Street
(C) CITY: Boston
(D) STATE: MA
(E) COUNTRY: USA
(F) ZIP: 02109

(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:
C--> **(A) APPLICATION NUMBER: US/08/887,505A**
C--> **(B) FILING DATE: 02-Jul-1997**
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: US 08/471,968
(B) FILING DATE: 06-JUN-1995

(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Kerner, Ann-Louise
(B) REGISTRATION NUMBER: 33,523
(C) REFERENCE/DOCKET NUMBER: 47508-250HYZ-040CIP

(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: (617) 526-6000
(B) TELEFAX: (617) 526-5000

(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid

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```
     59          (C) STRANDEDNESS: single
     60          (D) TOPOLOGY: linear
W--> 62     (ii) MOLECULE TYPE: DNA
     64    (iii) HYPOTHETICAL: NO
     66     (iv) ANTI-SENSE: YES
     68     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
     70 GGTGCACGGT CTACGAGACC                                          20
     72 (2) INFORMATION FOR SEQ ID NO: 2:
     74      (i) SEQUENCE CHARACTERISTICS:
     75          (A) LENGTH: 20 base pairs
     76          (B) TYPE: nucleic acid
     77          (C) STRANDEDNESS: single
     78          (D) TOPOLOGY: linear
W--> 80     (ii) MOLECULE TYPE: DNA
     82    (iii) HYPOTHETICAL: NO
     84     (iv) ANTI-SENSE: YES
     86     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
     88 CATGGTGCAC GGTCTACGAG                                          20
     90 (2) INFORMATION FOR SEQ ID NO: 3:
     92      (i) SEQUENCE CHARACTERISTICS:
     93          (A) LENGTH: 20 base pairs
     94          (B) TYPE: nucleic acid
     95          (C) STRANDEDNESS: single
     96          (D) TOPOLOGY: linear
W--> 98     (ii) MOLECULE TYPE: DNA
    100    (iii) HYPOTHETICAL: NO
    102     (iv) ANTI-SENSE: YES
    104     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
    106 GCTCATGGTG CACGGTCTAC                                          20
    109 (2) INFORMATION FOR SEQ ID NO: 4:
    111      (i) SEQUENCE CHARACTERISTICS:
    112          (A) LENGTH: 20 base pairs
    113          (B) TYPE: nucleic acid
    114          (C) STRANDEDNESS: single
    115          (D) TOPOLOGY: linear
W--> 117     (ii) MOLECULE TYPE: DNA
    119    (iii) HYPOTHETICAL: NO
    121     (iv) ANTI-SENSE: YES
    123     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
    125 GTGCTCATGG TGCACGGTCT                                          20
    127 (2) INFORMATION FOR SEQ ID NO: 5:
    129      (i) SEQUENCE CHARACTERISTICS:
    130          (A) LENGTH: 20 base pairs
    131          (B) TYPE: nucleic acid
    132          (C) STRANDEDNESS: single
    133          (D) TOPOLOGY: linear
W--> 134     (ii) MOLECULE TYPE: DNA
    136    (iii) HYPOTHETICAL: NO
    138     (iv) ANTI-SENSE: YES
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Input Set : A:\PTO.AMC.txt
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```
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
CGTGCTCATG GTGCACGGTC                                                  20
(2) INFORMATION FOR SEQ ID NO: 6:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 20 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
TTCGTGCTCA TGGTGCACGG                                                  20
(2) INFORMATION FOR SEQ ID NO: 7:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 20 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
GGATTCGTGC TCATGGTGCA                                                  20
(2) INFORMATION FOR SEQ ID NO: 8:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 20 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
TTAGGATTCG TGCTCATGGT                                                  20
(2) INFORMATION FOR SEQ ID NO: 9:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 20 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
GGTTTAGGAT TCGTGCTCAT                                                  20
(2) INFORMATION FOR SEQ ID NO: 10:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 20 base pairs
```

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```
     220           (B) TYPE: nucleic acid
     221           (C) STRANDEDNESS: single
     222           (D) TOPOLOGY: linear
W--> 224      (ii) MOLECULE TYPE: DNA
     226     (iii) HYPOTHETICAL: NO
     228      (iv) ANTI-SENSE: YES
     230      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
     232 TGAGGTTTAG GATTCGTGCT                                          20
     234 (2) INFORMATION FOR SEQ ID NO: 11:
     236       (i) SEQUENCE CHARACTERISTICS:
     237           (A) LENGTH: 20 base pairs
     238           (B) TYPE: nucleic acid
     239           (C) STRANDEDNESS: single
     240           (D) TOPOLOGY: linear
W--> 242      (ii) MOLECULE TYPE: DNA
     244     (iii) HYPOTHETICAL: NO
     246      (iv) ANTI-SENSE: YES
     248      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
     250 CTTTGAGGTT TAGGATTCGT                                          20
     252 (2) INFORMATION FOR SEQ ID NO: 12:
     254       (i) SEQUENCE CHARACTERISTICS:
     255           (A) LENGTH: 20 base pairs
     256           (B) TYPE: nucleic acid
     257           (C) STRANDEDNESS: single
     258           (D) TOPOLOGY: linear
W--> 260      (ii) MOLECULE TYPE: DNA
     262     (iii) HYPOTHETICAL: NO
     264      (iv) ANTI-SENSE: YES
     266      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
     268 TTCTTTGAGG TTTAGGATTC                                          20
     270 (2) INFORMATION FOR SEQ ID NO: 13:
     272       (i) SEQUENCE CHARACTERISTICS:
     273           (A) LENGTH: 20 base pairs
     274           (B) TYPE: nucleic acid
     275           (C) STRANDEDNESS: single
     276           (D) TOPOLOGY: linear
W--> 278      (ii) MOLECULE TYPE: DNA
     280     (iii) HYPOTHETICAL: NO
     282      (iv) ANTI-SENSE: YES
     284      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
     286 TACGTTTGGT TTTTCTTTGA                                          20
     288 (2) INFORMATION FOR SEQ ID NO: 14:
     290       (i) SEQUENCE CHARACTERISTICS:
     291           (A) LENGTH: 20 base pairs
     292           (B) TYPE: nucleic acid
     293           (C) STRANDEDNESS: single
     294           (D) TOPOLOGY: linear
W--> 296      (ii) MOLECULE TYPE: DNA
     298     (iii) HYPOTHETICAL: NO
```

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Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\09152003\H887505A.raw

```
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
GTTGGTGTTA CGTTTGGTTT                                                  20
(2) INFORMATION FOR SEQ ID NO: 15:
     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 20 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
GCACGACACT CATACTAACG                                                  20
(2) INFORMATION FOR SEQ ID NO: 16:
     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 20 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
GGCTGCACGA CACTCATACT                                                  20
(2) INFORMATION FOR SEQ ID NO: 17:
     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 20 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
TGGAGGCTGC ACGACACTCA                                                  20
(2) INFORMATION FOR SEQ ID NO: 18:
     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 20 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear
W--> (ii) MOLECULE TYPE: DNA
    (iii) HYPOTHETICAL: NO
     (iv) ANTI-SENSE: YES
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
GTCCTGGAGG CTGCACGACA                                                  20
(2) INFORMATION FOR SEQ ID NO: 19:
     (i) SEQUENCE CHARACTERISTICS:
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VERIFICATION SUMMARY DATE: 09/15/2003
PATENT APPLICATION: **US/08/887,505A** TIME: 10:39:54

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\09152003\H887505A.raw**

L:35 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:36 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:62 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:80 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:98 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:117 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4
L:134 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5
L:152 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6
L:170 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7
L:188 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8
L:206 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9
L:224 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10
L:242 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11
L:260 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12
L:278 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13
L:296 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14
L:314 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15
L:332 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16
L:350 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17
L:368 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=18
L:386 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=19
L:404 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=20
L:422 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=21
L:440 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22
L:458 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=23
L:476 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=24
L:494 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25
L:512 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=26
L:530 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27
L:548 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28
L:566 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29
L:584 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30
L:602 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31
L:620 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32
L:638 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33
L:656 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34
L:674 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35
L:692 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:710 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:728 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38
L:746 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39
L:764 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40
L:782 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41
L:800 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42
L:818 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=43
L:836 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44
L:854 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=45
L:872 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=46

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Input Set : A:\PTO.AMC.txt
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L:890 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=47
L:908 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=48
L:926 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=49
L:944 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50